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Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: Source: Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY of,
- TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS: http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual - ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06





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RAW SEQUENCE LISTING PATENT APPLICATION: US/10/528,260A

DATE: 08/14/2006 TIME: 13:29:48

Input Set : E:\P1983R1.txt

Output Set: N:\CRF4\08142006\J528260A.raw

3 <110> APPLICANT: HENRY CHIU HILARY CLARK KATHRYN DENNIS SHERMAN FONG JILL R. SCHOENFELD WILLIAM I. WOOD

11 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING IMMUNE

DISORDERS

13 <130> PILE REFERENCE: P1983R1 15 <140> CURRENT APPLICATION NUMBER: US 10/528,260A 16 <141> CURRENT FILING DATE: 2005-03-16

18 <150> PRIOR APPLICATION NUMBER: US 60/411,392 19 <151> PRIOR FILING DATE: 2002-09-16

21 <150> PRIOR APPLICATION NUMBER: US03/029097

22 <151> PRIOR FILING DATE: 2003-09-15

24 <150> NUMBER OF SEQ ID NOS 506

Does Not Comply Corrected Diskette Needed

BRRORED SEQUENCES

B--> 17766 <210> SBQ ID NO: 177 E--> 23561 <210> SEQ ID NO: 225 B--> 23579 <210> SBQ ID NO: 227 B--> 23961 <210> SEQ ID NO: 235 E--> 24152 <210> SEQ ID NO: 239 E--> 24938 <210> SEQ ID NO: 247 E--> 25002 <210> SEQ ID NO: 249 E--> 25335 <210> SEQ ID NO: 253 E--> 34738 <210> BEQ ID NO: 331 E--> 34773 <210> SEQ ID NO: 333 E--> 35071 <210> SEQ ID NO: 339 E--> 35313 <210> SEQ ID NO: 343 B--> 39779 <210> SEQ ID NO: 387 E--> 41386 <210> 5EQ ID NO: 405 R--> 42010 <210> SEQ ID NO: 413 E--> 42381 <210> SEQ ID NO: 417 B--> 42426 <210> SEQ ID NO: 419 B--> 42897 <210> SEQ ID NO: 425 B--> 42917 <210> SEQ. ID, NO: 427

B--> 42975 <210> SEQ ID NO: 429



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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/528,260A DATE: 08/14/2006 TIME: 13:29:51

Input Set : B:\P1983R1.txt

Output Set: N:\CRF4\081420D6\J528260A.raw

Skipped Sequences (NEW RULES):
Sequence (s) missing. If intentional, please use the following format for

each skipped sequence. <210> sequence id number

<400> sequence id number

of See p-4 fr Sample

Seq#:176,224,226,234,238,246,248,252,330,332,338,342,386,404,412,416,418,424 Seq#:426,428,430,480,492,494

the lbore sequence are missing

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VERIFICATION SUMMARY PATENT APPLICATION: US/10/528,260A

DATE: 08/14/2006 TIME: 13:29:51

Imput Set : R:\P1983R1.txt

Output Set: N:\CRF4\08142006\J528260A.raw

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L:17766 M:214 B: (33) Seq.# missing, SPQ ID NO:176
    L:20650 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:200 after pos.:15
    L:22238 M:341 W: (46) "n" or "Xaa" used, for SEQ 1D#:213 after pos.:1350
    L:23561 M:214 B: (33) Beq.# missing, SEQ ID NO:224
    L:23579 M:214 B: (33) Seq.# missing. SEQ ID NO:226
    L:23599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:227 after pos.:100
    M:341 Repeated in SeqNo=227
    L:23961 M:214 B: (33) Seq.# missing, SEQ ID NO:234
L:24152 M:214 B: (33) Seq.# missing, SEQ ID NO:238
    L:24194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:239 after pos.:400
    M:341 Repeated in SeqNo=239
    L:24938 M:214 E: (33) Seq.# missing, SEQ ID NO:246
    L:25002 M:214 R: (33) Seq. # missing, 880 ID NO:248
     L:25335 M:214 B: (33) Seq.# missing, SEQ ID NO:252
     L:27797 M:341 W: (46) "n° or "Xaa" used, for SEQ ID#:281 after pos.:2500
     L:33378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:321 after pos.:2700
L:33599 N:341 W: (46) "n" or "Xaa" used, for SEQ ID#:322 after pos.:900
     L:34738 M:214 B: (33) Seq.# missing, SEQ ID NO:330
     L:34753 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:331 after pos.:100
     L:34773 M:214 B: (33) Seq.# missing, SEQ ID NO:332
   L:35071 M:214 E: (33) Seq.# missing, SEQ ID NO:338
     L:35313 M:214 E: (33) Seq.# missing, SEQ ID NO:342
L:39779 M:214 E: (33) Seq.# missing, SEQ ID NO:386
     L:40746 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:396 after pos.:315
   L:41386 M:214 B: (33) Seq.# missing, SEQ ID NO:404
     L:42010 M:214 B: (33) Seq.# missing, SEQ ID NO:412
     L:42355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:415 after pos.:0
 L:42381 M:214 R: (33) Seq.# missing, SEQ ID NO:416
... ; L:42422 M:341 W: (46) *n" or "Xaa" used, for SEQ ID#:417 after pos.:750
     L:42426 M:214 B: (33) Seq.# missing, SEQ ID NO:418
 L:42861 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:423 after pos.:0
 M:341 Repeated in SeqNo=423
     L:42897 M:214 B: (33) Seq.# missing, SEQ ID NO:424
     L:42917 M:214 E: (33) Seq.# missing, SEQ ID NO:426
 L:42975 M:214 B: (33) Seq.# missing, SEQ 1D NO:428
     L:43009 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:429 after pos.:450
  L:43013 M:214 B: (33) Seq.# missing, SEQ ID NO:430
  L:49277 M:214 B: (33) Seq.# missing, SEQ ID NO:480
     L:50541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:491 after pos.:1950
   M:341 Repeated in SeqNo=491
  L:50645 M:214 B: (33); Seq.# missing, SEQ ID NO:492
L:50859 M:214 E: (33); Seq.# missing, SEQ ID NO:494
L:50875 M:341 W: (46) *n* or "Kaa* used, for SEQ ID#:495 after pos.:0
 M:341 Repeated in SeqNo-495

L:51187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:496 after pos.:0

M:341 Repeated in SeqNo-496
- .... M:341 Repeated in SeqNo=495
```

BEST AVAILABLE COPY 10/528, 260H

(end of Sequence 175)

Sar Arg Phe Lys Lys Arg Met Leu Ile Glu Gln Leu Glu Asn Phe

Leu Asp Glu Ile His Arg Arg Ala Asn Gln Ile Asn His Ile Asn

where is Sequence 176?

<210> 177

<211> 2533

<212> DNA

<213> Homo sapiens

<400> 177

ccaageccat gagggeegeg egeceggeeg eeggtgetga egagaeggag 50

(Sample of numerous missing Sequence (Sting file).